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measured from N terminal end are both deleted, and at least one amino acid residue from  $111^{th}$  amino acid to  $128^{th}$  amino acid residues or at least one amino acid residue from  $131^{st}$  amino acid to  $133^{rd}$  amino acid residues as measured from N terminal end is deleted.

- 3. (Four times Amended) An isolated polypeptide having an amino acid sequence of natural human Fas ligand (SEQ ID NO:17) wherein all of the  $8^{th}$  amino acid to  $69^{th}$  amino acid residues as measured from N terminal end are deleted,  $129^{th}$  amino acid and  $130^{th}$  amino acid residues as measured from N terminal end are both deleted, and at least one amino acid residue from  $111^{th}$  amino acid to  $128^{th}$  amino acid residues or at least one amino acid residues from  $131^{st}$  amino acid to  $133^{rd}$  amino acid residues as measured from N terminal end is deleted.
- 5. (Twice Amended) An isolated DNA coding for the polypeptide of claim 2.
- 6. (Thrice Amended) A soluble Fas ligand which inhibits Fasmediated apoptosis and which comprises the amino acid sequence represented from Gln of the  $130^{\rm th}$  amino acid to C terminal amino acid residue as measured from N-terminal end of natural human Fas ligand (SEQ ID NO:17).

- 8. (Amended) An isolated DNA coding for the polypeptide of claim 3.
- 9. (Amended) An isolated DNA coding for the polypeptide of claim 4.
- 10. (Four times amended) An isolated polypeptide having an amino acid sequence of natural human Fas ligand (SEQ ID NO:17) wherein the 129<sup>th</sup> amino acid and 130<sup>th</sup> amino acid residues as measured from N terminal end are both deleted, and at least one amino acid residue from 111<sup>th</sup> amino acid to 128<sup>th</sup> amino acid residues or at least one amino acid residue from 131<sup>st</sup> amino acid to 133<sup>rd</sup> amino acid residues as measured from N terminal end is deleted, wherein said polypeptide has membrane binding activity and induces Fas-mediated apoptotic activity.
- 11. (Four times amended) An isolated polypeptide having an amino acid sequence of natural human Fas ligand (SEQ ID NO:17) wherein all of the  $8^{\rm th}$  amino acid to  $69^{\rm th}$  amino acid residues as measured from N terminal end are deleted,  $129^{\rm th}$  amino acid and  $130^{\rm th}$  amino acid residues as measured from N terminal end are both deleted, and at least one amino acid residue from  $111^{\rm th}$  amino acid to  $128^{\rm th}$  amino acid residues or at least one amino acid residues from  $131^{\rm st}$  amino acid to  $133^{\rm rd}$  amino acid residues as measured from N terminal end is deleted, wherein said

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polypeptide has membrane binding activity and induces Fasmediated apoptotic activity.

12. (Amended) An isolated peptide having an amino acid sequence of natural human Fas ligand (SEQ ID NO:17) wherein at least four amino acid residues, including 128<sup>th</sup> and 131<sup>st</sup> amino acid residues are continuously deleted from the 111<sup>th</sup> amino acid to the 133<sup>rd</sup> amino acid residues as measured from N terminal end. wherein the 129<sup>th</sup> amino acid and 130<sup>th</sup> amino acid residues as measured from N terminal end are both deleted, and at least one amino acid residue from 111<sup>th</sup> amino acid to 128<sup>th</sup> amino acid residues or at least one amino acid residue from 131<sup>st</sup> amino acid to 133<sup>rd</sup> amino acid residues as measured from N terminal end is deleted.